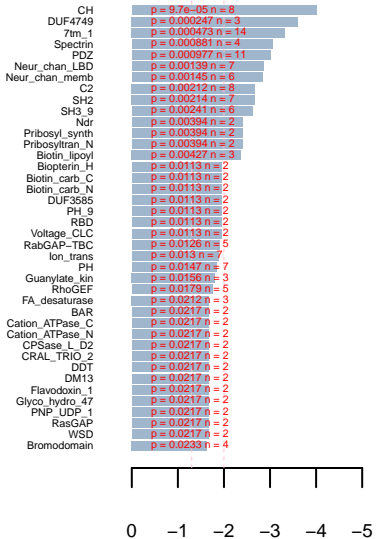


## OeCa\_F domain



log(p)

n = 672 genes | p adjust BH FALSE